==========

Sequence Listing could not be accepted due to errors.
See attached Validation Report.
If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).
Reviewer: Anne Corrigan
Timestamp: Thu Jun 07 20:19:22 EDT 2007

==========

****

Reviewer Comments:
<210> 1
<211> 23
<212> DNA
<213> Artificial Sequence
<220>
<223> CodY target sequence

Please explain the source of "<223> CodY target sequence."

<210> 15
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> consensus sequence GTP binding motif in CodY homologs
<220>
<221> Xaa
<222> (2)..(2)
<223> Arg
<220>
<221> Xaa
<222> (3)..(3)
<223> Met, Gly, Ile, Lys, Gln
<220>
<221> Xaa
<222> (4)..(4)
<223> Gly
<220>

<221> Xaa
<222> (5)..(5)
<223> Thr
<220>
<221> Xaa
<222> (7)..(7)
<223> Ser
<400> 15
Gly Gly Glu Arg Leu Gly Thr Thr
  1   5

There are no Xaa's at locations 2-4, and 5-7. Why are there Xaa explanations? e.g., "Gly" is at location 2--it can only represent itself; "Glu" is at location 3. Same type of error in Sequence 15.

<220>
<221> Variant
<222> (2)..(2)
<223> Ala, Lys
<220>
<221> Variant
<222> (3)..(3)
<223> Phe, Ile
<400> 24
Asp Arg Val Gly
  1

The <222> and <223> responses above are invalid. "Arg" is at location 2--it can only represent itself. If you want location 2 to represent Arg, Ala, or Lys, please use "Xaa" instead of "Arg." Same problem with the <222> (3)..(3) and the <223> line below it.

Same type of error in Sequence 29.

 <210> 226
<211> 30
<212> DNA
<213> Artificial Sequence
<220>
<223> MUT16

The <223> response above is an insufficient explanation for <213>

Artificial Sequence. Please give the source of "MUT16." Same type of error throughout the submitted file.
****

Validated By CRFValidator v 1.0.2

Application No: 10562601 Version No: 1.0

Input Set:

Output Set:

Started: 2007-06-07 15:32:33.514
Finished: 2007-06-07 15:32:36.262
Elapsed: 0 hr(s) 0 min(s) 2 sec(s) 748 ms
Total Warnings: 207
Total Errors: 12
No. of SeqIDs Defined: 234
Actual SeqID Count: 234

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W	213	Artificial or Unknown found in <213> in SEQ ID (2)
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Input Set:

Output Set:

Started: 2007-06-07 15:32:33.514
Finished: 2007-06-07 15:32:36.262
Elapsed: 0 hr(s) 0 min(s) 2 sec(s) 748 ms
Total Warnings: 207
Total Errors: 12
No. of SeqIDs Defined: 234
Actual SeqID Count: 234

Error code		Error Description
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W	213	Artificial or Unknown found in <213> in SEQ ID (40)
W	213	Artificial or Unknown found in <213> in SEQ ID (41) This error has occured more than 20 times, will not be displayed
E	257	Invalid sequence data feature in <221> in SEQ ID (214)
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SEQUENCE LISTING

<110> Hengst den, Christiaan D.
Gajic, Olivera
Kuipers, Oscar P.
Kok, Jan
Sikkema, Jan
Geurts, Johannes M.W.
Nauta, Arjen

<120> Methods and means for regulating gene expression

<130> P63590US00

<140> 10562601
<141> 2007-06-07

<150> US 10/562,601
<151> 2005-12-28

<150> PCT/NL2004/000474
<151> 2004-07-02

<150> EP 03077074.7
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<210> 3
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<223> primer opp1

<400> 3
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<223> primer opp2

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<210> 5
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<400> 5
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<400> 14
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<220>
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<400> 16
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<210> 17
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  1   5

<210> 18
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<213> Bacillus halodurans

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  1   5

<210> 19

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<400> 19
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  1   5

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<400> 20
Asn Arg Glu Arg Leu Gly Thr Leu
  1   5

<210> 21
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<220>
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  1   5

<210> 22
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<220>
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Asp Ala Phe Gly
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<210> 26

<211> 4

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<220>
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<211> 4
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<220>
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<400> 27
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  1

<210> 28
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<210> 29
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<220>
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<220>

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<210> 30
<211> 4
<212> PRT
<213> Escherichia coli

<220>
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<211> 4
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<223> /note="Putative GTP binding motif G4"

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<210> 32
<211> 4
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<213> Bacillus stearothermophilus

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<222> (1)..(4)
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Ile Leu Asn Asp
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<220>
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<210> 36
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<212> PRT
<213> Enterococcus faecalis

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<400> 36
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<210> 37
<211> 4
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<400> 37
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<210> 38
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<212> PRT
<213> Lactococcus lactis

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<400> 38
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<210> 39
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<210> 41
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<223> presence of the putative CodY box in optA

<400> 42
aaattttctg acaataataa aaattg 26

<210> 43
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<210> 62
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<210> 63
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<210> 72
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<210> 73
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<220>
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attttcaga aaaat 15

<210> 74
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<212> DNA
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